



1600

RAW SEQUENCE LISTING

DATE: 03/29/2004

PATENT APPLICATION: US/10/019,114A

TIME: 09:56:27

Input Set : A:\3-19-04 KATO revised sequence listing.txt

Output Set: N:\CRF4\03292004\J019114A.raw

5 <110> APPLICANT: Seishi KATO
 6 Tomoko KIMURA
 9 <120> TITLE OF INVENTION: Human proteins having hydrophobic domains and DNAs encoding
 these
 11 proteins
 14 <130> FILE REFERENCE: 2003-0986/WMC/01791
 17 <140> CURRENT APPLICATION NUMBER: 10/019,114A
 C--> 18 <141> CURRENT FILING DATE: 2001-12-21
 21 <150> PRIOR APPLICATION NUMBER: JP 11-188835
 22 <151> PRIOR FILING DATE: 1999-07-02
 25 <160> NUMBER OF SEQ ID NOS: 30
 28 <210> SEQ ID NO: 1
 29 <211> LENGTH: 233
 30 <212> TYPE: PRT
 31 <213> ORGANISM: Homo sapiens
 35 <400> SEQUENCE: 1
 37 Met Trp Gln Leu Leu Ala Ala Ala Cys Trp Met Leu Leu Leu Gly Ser
 39 1 5 10 15
 41 Met Tyr Gly Tyr Asp Lys Lys Gly Asn Asn Ala Asn Pro Glu Ala Asn
 43 20 25 30
 45 Met Asn Ile Ser Gln Ile Ile Ser Tyr Trp Gly Tyr Pro Tyr Glu Glu
 47 35 40 45
 49 Tyr Asp Val Thr Thr Lys Asp Gly Tyr Ile Leu Gly Ile Tyr Arg Ile
 51 50 55 60
 53 Pro His Gly Arg Gly Cys Pro Gly Arg Thr Ala Pro Lys Pro Ala Val
 55 65 70 75 80
 57 Tyr Leu Gln His Gly Leu Ile Ala Ser Ala Ser Asn Trp Ile Cys Asn
 59 85 90 95
 61 Leu Pro Asn Asn Ser Leu Ala Phe Leu Leu Ala Asp Ser Gly Tyr Asp
 63 100 105 110
 65 Val Trp Leu Gly Asn Ser Arg Gly Asn Thr Trp Ser Arg Lys His Leu
 67 115 120 125
 69 Lys Leu Ser Pro Lys Ser Pro Glu Tyr Trp Ala Phe Ser Leu Asp Glu
 71 130 135 140
 73 Met Ala Lys Tyr Asp Leu Pro Ala Thr Ile Asn Phe Ile Ile Glu Lys
 75 145 150 155 160
 77 Thr Gly Gln Lys Arg Leu Tyr Tyr Val Gly His Ser Gln Gly Thr Thr
 79 165 170 175
 81 Ile Ala Phe Ile Ala Phe Ser Thr Asn Pro Glu Leu Ala Lys Lys Ile
 83 180 185 190
 85 Lys Ile Phe Phe Ala Leu Ala Pro Val Val Thr Val Lys Tyr Thr Gln
 87 195 200 205
 89 Ser Pro Met Lys Lys Leu Thr Leu Ser Arg Arg Val Val Lys Val
 91 210 215 220

ENTERED

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93 Cys Asp Phe Pro Ser Phe Asn Leu Lys
95 225                230
98 <210> SEQ ID NO: 2
99 <211> LENGTH: 273
100 <212> TYPE: PRT
101 <213> ORGANISM: Homo sapiens
103 <400> SEQUENCE: 2
105 Met Arg Gly Ser Gln Glu Val Leu Leu Met Trp Leu Leu Val Leu Ala
107 1                5                10                15
109 Val Gly Gly Thr Glu His Ala Tyr Arg Pro Gly Arg Arg Val Cys Ala
111 20                25                30
113 Val Arg Ala His Gly Asp Pro Val Ser Glu Ser Phe Val Gln Arg Val
115 35                40                45
117 Tyr Gln Pro Phe Leu Thr Thr Cys Asp Gly His Arg Ala Cys Ser Thr
119 50                55                60
121 Tyr Arg Thr Ile Tyr Arg Thr Ala Tyr Arg Arg Ser Pro Gly Leu Ala
123 65                70                75                80
125 Pro Ala Arg Pro Arg Tyr Ala Cys Cys Pro Gly Trp Lys Arg Thr Ser
127 85                90                95
129 Gly Leu Pro Gly Ala Cys Gly Ala Ala Ile Cys Gln Pro Pro Cys Arg
131 100               105               110
133 Asn Gly Gly Ser Cys Val Gln Pro Gly Arg Cys Arg Cys Pro Ala Gly
135 115               120               125
137 Trp Arg Gly Asp Thr Cys Gln Ser Asp Val Asp Glu Cys Ser Ala Arg
139 130               135               140
141 Arg Gly Gly Cys Pro Gln Arg Cys Val Asn Thr Ala Gly Ser Tyr Trp
143 145               150               155               160
145 Cys Gln Cys Trp Glu Gly His Ser Leu Ser Ala Asp Gly Thr Leu Cys
147 165               170               175
149 Val Pro Lys Gly Gly Pro Pro Arg Val Ala Pro Asn Pro Thr Gly Val
151 180               185               190
153 Asp Ser Ala Met Lys Glu Glu Val Gln Arg Leu Gln Ser Arg Val Asp
155 195               200               205
157 Leu Leu Glu Glu Lys Leu Gln Leu Val Leu Ala Pro Leu His Ser Leu
159 210               215               220
161 Ala Ser Gln Ala Leu Glu His Gly Leu Pro Asp Pro Gly Ser Leu Leu
163 225               230               235               240
165 Val His Ser Phe Gln Gln Leu Gly Arg Ile Asp Ser Leu Ser Glu Gln
167 245               250               255
169 Ile Ser Phe Leu Glu Glu Gln Leu Gly Ser Cys Ser Cys Lys Lys Asp
171 260               265               270
173 Ser
175 <210> SEQ ID NO: 3
176 <211> LENGTH: 282
177 <212> TYPE: PRT
178 <213> ORGANISM: Homo sapiens
180 <400> SEQUENCE: 3
182 Met Ser Gly Ser Ser Leu Pro Ser Ala Leu Ala Leu Ser Leu Leu Leu
184 1                5                10                15

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```

186 Val Ser Gly Ser Leu Leu Pro Gly Pro Gly Ala Ala Gln Asn Glu Pro
188      20      25      30
190 Arg Ile Val Thr Ser Glu Glu Val Ile Ile Arg Asp Ser Pro Val Leu
192      35      40      45
194 Pro Val Thr Leu Gln Cys Asn Leu Thr Ser Ser Ser His Thr Leu Thr
196      50      55      60
198 Tyr Ser Tyr Trp Thr Lys Asn Gly Val Glu Leu Ser Ala Thr Arg Lys
200      65      70      75      80
202 Asn Ala Ser Asn Met Glu Tyr Arg Ile Asn Lys Pro Arg Ala Glu Asp
204      85      90      95
206 Ser Gly Glu Tyr His Cys Val Tyr His Phe Val Ser Ala Pro Lys Ala
208      100      105      110
210 Asn Ala Thr Ile Glu Val Lys Ala Ala Pro Asp Ile Thr Gly His Lys
212      115      120      125
214 Arg Ser Glu Asn Lys Asn Glu Gly Gln Asp Ala Thr Met Tyr Cys Lys
216      130      135      140
218 Ser Val Gly Tyr Pro His Pro Asp Trp Ile Trp Arg Lys Lys Glu Asn
220      145      150      155      160
222 Gly Met Pro Met Asp Ile Val Asn Thr Ser Gly Arg Phe Phe Ile Ile
224      165      170      175
226 Asn Lys Glu Asn Tyr Thr Glu Leu Asn Ile Val Asn Leu Gln Ile Thr
228      180      185      190
230 Glu Asp Pro Gly Glu Tyr Glu Cys Asn Ala Thr Asn Ala Ile Gly Ser
232      195      200      205
234 Ala Ser Val Val Thr Val Leu Arg Val Arg Ser His Leu Ala Pro Leu
236      210      215      220
238 Trp Pro Phe Leu Gly Ile Leu Ala Glu Ile Ile Ile Leu Val Val Ile
240      225      230      235      240
242 Ile Val Val Tyr Glu Lys Arg Lys Arg Pro Asp Glu Val Pro Asp Asp
244      245      250      255
246 Asp Glu Pro Ala Gly Pro Met Lys Thr Asn Ser Thr Asn Asn His Lys
248      260      265      270
250 Asp Lys Asn Leu Arg Gln Arg Asn Thr Asn
252      275      280
255 <210> SEQ ID NO: 4
256 <211> LENGTH: 238
257 <212> TYPE: PRT
258 <213> ORGANISM: Homo sapiens
260 <400> SEQUENCE: 4
262 Met Ser Leu Asn Glu His Ser Met Gln Ala Leu Ser Trp Arg Lys Leu
264      1      5      10      15
266 Tyr Leu Ser Arg Ala Lys Leu Lys Ala Ser Ser Arg Thr Ser Ala Leu
268      20      25      30
270 Leu Ser Gly Phe Ala Met Val Ala Met Val Glu Val Gln Leu Asp Ala
272      35      40      45
274 Asp His Asp Tyr Pro Pro Gly Leu Leu Ile Ala Phe Ser Ala Cys Thr
276      50      55      60
278 Thr Val Leu Val Ala Val His Leu Phe Ala Leu Met Ile Ser Thr Cys
280      65      70      75      80

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282 Ile Leu Pro Asn Ile Glu Ala Val Ser Asn Val His Asn Leu Asn Ser
284      85      90      95
286 Val Lys Glu Ser Pro His Glu Arg Met His Arg His Ile Glu Leu Ala
288      100      105      110
290 Trp Ala Phe Ser Thr Val Ile Gly Thr Leu Leu Phe Leu Ala Glu Val
292      115      120      125
294 Val Leu Leu Cys Trp Val Lys Phe Leu Pro Leu Lys Lys Gln Pro Gly
296      130      135      140
298 Gln Pro Arg Pro Thr Ser Lys Pro Pro Ala Ser Gly Ala Ala Ala Asn
300 145      150      155      160
302 Val Ser Thr Ser Gly Ile Thr Pro Gly Gln Ala Ala Ala Ile Ala Ser
304      165      170      175
306 Thr Thr Ile Met Val Pro Phe Gly Leu Ile Phe Ile Val Phe Ala Val
308      180      185      190
310 His Phe Tyr Arg Ser Leu Val Ser His Lys Thr Asp Arg Gln Phe Gln
312      195      200      205
314 Glu Leu Asn Glu Leu Ala Glu Phe Ala Arg Leu Gln Asp Gln Leu Asp
316      210      215      220
318 His Arg Gly Asp His Pro Leu Thr Pro Gly Ser His Tyr Ala
320 225      230      235
324 <210> SEQ ID NO: 5
325 <211> LENGTH: 372
326 <212> TYPE: PRT
327 <213> ORGANISM: Homo sapiens
329 <400> SEQUENCE: 5
331 Met Leu Ala Asn Ser Ser Ser Thr Asn Ser Ser Val Leu Pro Cys Pro
333 1      5      10      15
335 Asp Tyr Arg Pro Thr His Arg Leu His Leu Val Val Tyr Ser Leu Val
337      20      25      30
339 Leu Ala Ala Gly Leu Pro Leu Asn Ala Leu Ala Leu Trp Val Phe Leu
341      35      40      45
343 Arg Ala Leu Arg Val His Ser Val Val Ser Val Tyr Met Cys Asn Leu
345      50      55      60
347 Ala Ala Ser Asp Leu Leu Phe Thr Leu Ser Leu Pro Val Arg Leu Ser
349 65      70      75      80
351 Tyr Tyr Ala Leu His His Trp Pro Phe Pro Asp Leu Leu Cys Gln Thr
353      85      90      95
355 Thr Gly Ala Ile Phe Gln Met Asn Met Tyr Gly Ser Cys Ile Phe Leu
357      100      105      110
359 Met Leu Ile Asn Val Asp Arg Tyr Ala Ala Ile Val His Pro Leu Arg
361      115      120      125
363 Leu Arg His Leu Arg Arg Pro Arg Val Ala Arg Leu Leu Cys Leu Gly
365      130      135      140
367 Val Trp Ala Leu Ile Leu Val Phe Ala Val Pro Ala Ala Arg Val His
369 145      150      155      160
371 Arg Pro Ser Arg Cys Arg Tyr Arg Asp Leu Glu Val Arg Leu Cys Phe
373      165      170      175
375 Glu Ser Phe Ser Asp Glu Leu Trp Lys Gly Arg Leu Leu Pro Leu Val
377      180      185      190

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```

379 Leu Leu Ala Glu Ala Leu Gly Phe Leu Leu Pro Leu Ala Ala Val Val
381      195      200      205
383 Tyr Ser Ser Gly Arg Val Phe Trp Thr Leu Ala Arg Pro Asp Ala Thr
385      210      215      220
387 Gln Ser Gln Arg Arg Arg Lys Thr Val Arg Leu Leu Leu Ala Asn Leu
389 225      230      235      240
391 Val Ile Phe Leu Leu Cys Phe Val Pro Tyr Asn Ser Thr Leu Ala Val
393      245      250      255
395 Tyr Gly Leu Leu Arg Ser Lys Leu Val Ala Ala Ser Val Pro Ala Arg
397      260      265      270
399 Asp Arg Val Arg Gly Val Leu Met Val Met Val Leu Leu Ala Gly Ala
401      275      280      285
403 Asn Cys Val Leu Asp Pro Leu Val Tyr Tyr Phe Ser Ala Glu Gly Phe
405      290      295      300
407 Arg Asn Thr Leu Arg Gly Leu Gly Thr Pro His Arg Ala Arg Thr Ser
409 305      310      315      320
411 Ala Thr Asn Gly Thr Arg Ala Ala Leu Ala Gln Ser Glu Arg Ser Ala
413      325      330      335
415 Val Thr Thr Asp Ala Thr Arg Pro Asp Ala Ala Ser Gln Gly Leu Leu
417      340      345      350
419 Arg Pro Ser Asp Ser His Ser Leu Ser Ser Phe Thr Gln Cys Pro Gln
421      355      360      365
423 Asp Ser Ala Leu
425      370
429 <210> SEQ ID NO: 6
430 <211> LENGTH: 146
431 <212> TYPE: PRT
432 <213> ORGANISM: Homo sapiens
434 <400> SEQUENCE: 6
436 Met Trp Lys Leu Trp Arg Ala Glu Glu Gly Ala Ala Ala Leu Gly Gly
438 1      5      10      15
440 Ala Leu Phe Leu Leu Phe Ala Leu Gly Val Arg Gln Leu Leu Lys
442      20      25      30
444 Gln Arg Arg Pro Met Gly Phe Pro Pro Gly Pro Pro Gly Leu Pro Phe
446      35      40      45
448 Ile Gly Asn Ile Tyr Ser Leu Ala Ala Ser Ser Glu Leu Pro His Val
450      50      55      60
452 Tyr Met Arg Lys Gln Ser Gln Val Tyr Gly Glu Val Gln Pro Arg Arg
454 65      70      75      80
456 Ala Pro Gly Arg Glu Gly Arg Gln Ala Gly Pro Gly Trp Pro Gly Pro
458      85      90      95
460 Ser Trp Leu Asp Leu Trp Pro Pro Leu Gly Arg Leu Val Gly Thr Ser
462      100      105      110
464 Pro Cys Ala Gly Cys Pro Leu Arg Asp Thr Arg Phe Pro Gly Leu Glu
467      115      120      125
469 Gly Arg Ser Pro Arg Arg Arg Ala Pro Leu Gln Gly Glu Pro Arg Pro
471      130      135      140
473 Cys Arg
475 145

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VERIFICATION SUMMARY

DATE: 03/29/2004

PATENT APPLICATION: US/10/019,114A

TIME: 09:56:28

Input Set : A:\3-19-04 KATO revised sequence listing.txt

Output Set: N:\CRF4\03292004\J019114A.raw

L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date